

0100  
23-01 0280

OIPE

ENTERED

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/817,360

DATE: 04/05/2001  
TIME: 12:14:04

Input Set : A:\SeqList for UCSF-129CIP.txt  
Output Set : N:\CRF3\04052001\I817360.raw

4 <110> APPLICANT: German, Michael S.  
5 Lin, Joseph  
7 <120> TITLE OF INVENTION: PRODUCTION OF PANCREATIC ISLET CELLS  
8 AND DELIVERY OF INSULIN  
13 <130> FILE REFERENCE: UCSF-129CIP  
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C--> 15 <141> CURRENT FILING DATE: 2001-03-20  
15 <150> PRIOR APPLICATION NUMBER: 09/535,145  
16 <151> PRIOR FILING DATE: 2000-03-24  
18 <150> PRIOR APPLICATION NUMBER: 60/128,180  
19 <151> PRIOR FILING DATE: 1999-04-06  
21 <160> NUMBER OF SEQ ID NOS: 19  
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
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27 <212> TYPE: DNA  
28 <213> ORGANISM: Homo Sapiens  
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33 tctaacttgc agttaataa atcaggcaag gctggcctat gaggcagaca agtgtaaga 180  
34 aggagaagga ggaggagaag gagaaggaga aagaagaaga aggaggagaa gaagaagaag 240  
35 aagaagaaga agaaggagg gaggaggagg aggaggagga agcagcagca gcagcagcag 300  
36 cttgaatgga cagtgttcc ccttgcctag aaaatgggac cattatttct tttctaattct 360  
37 gacccccaga ctcaggactt cctctatttt ctgcattttg ggtctctttg ttttgccttg 420  
38 aaaaaaaatg ttttctccca aatcaaggag cagtagctgg tgcaaggga aatctagggc 480  
39 taggagtctt aagatatgac ttctatgtgg ttctgataga acttgctggg tgaccttgag 540  
40 agagtcactc cccctctctg ggcttgatt tttcatctt taaagaaggc ctcaaattcc 600  
41 cattcttatg agaagaagac aagctcctag tgagtgggta cctaaggag cagctgcagc 660  
42 aaaatgctaa cctgacagtc ccagatggtc cctttattgg ttctgacct ggtctcaggc 720  
43 ttcattttcc cacagcaagg gaaggagcct gctcacagag caccagctaa gatcagcagg 780  
44 accgcgccac acccccgccc agtcctagag ccccccctct gctggttcct gagcatacca 840  
45 ccctcttctt tggaggaaaa ttgccccca agcagcctag gcggaagag gctatcacta 900  
46 gggcagactc acagacctac ctcatccct caccacccc tacagtctcg aagtcgggtc 960  
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59 cagtgaactg tgattgcacc actgcagctc agcctgggag acagaaggag accgtttttt 1740

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62	gggatactat	ctacaagctg	tagtaggctt	gtagtaatgg	aatgtccgct	tgagggggtcc	1920
63	ccgcacagcc	aacccccgcc	tctggagtg	gatctatggg	ggtgggggtc	taagcgctc	1980
64	tggggagtg	gaggtagcat	ctcaggggtg	ggcagaggct	cggacacccc	caaaagggtct	2040
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66	agagaggcag	tgaacacca	ggagagcagg	gcgtccttta	gaattcctgg	acccttctcc	2160
67	aggctgctag	tcaggacaat	gagctcgtgg	ttgtctttgc	cactatcttc	ctgtgcgatt	2220
68	tcagacaagc	cacctccctc	actaagccta	aatttcccc	tgtgtaacgt	gcaggcattg	2280
69	taccgtāgag	gcattcaagt	cccctccagt	acagatgcta	aggaagata	ggctaggagc	2340
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71	tttgaggaac	cgagagttgc	tgggacccag	cccgcctcg	agagagcaaa	cagagcggcg	2460
72	ctccctcccc	ccgaccccg	ccctttgtcc	ggaatccagc	tgtgctgcg	gggaggagcg	2520
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103	cgaactgcgc	ctgaaacgtg	agctgcgctg	caggtgcctg	gagcacccgc	catctttttt	4380
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105	tatatggccc	ctccctgttt	ttacacattt	gtatttatta	atgagatttc	acagcaggga	4500
106	aaagcctata	ttttggatat	tagattattt	agggattgct	ggatgacatt	taagccaata	4560
107	aaaaaaaaatg	gaccttcaag	aagccttggc	aagatgactc	cattgtgtgt	tggggagagg	4620
108	agggccacag	tcactacagc	tgagggaag	cacttctgtc	caaagagagg	gatgacactc	4680

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111 gagtgggtta gaagaaagac aagcctccaa ctaggacaac tgactctcac ttgctggccc 4860
112 tttccccaac tccaccagcc tagctttaga gcaactgttg gttgcacttg gggaagggat 4920
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116 gagcgtcttg ctggaatata cagcaacgcc ccctccctca tcacctggca gccttgattg 5160
117 aaaacttatt aagaactgt tcaaggtttc cagccacacc atgtctctta ctggcaagggt 5220
118 ggaataggac tgggtcagca tgagcactga aatctgtccc aggagtgccca gtagagcacc 5280
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120 &lt;210&gt; SEQ ID NO: 2

121 &lt;211&gt; LENGTH: 214

122 &lt;212&gt; TYPE: PRT

123 &lt;213&gt; ORGANISM: Homo Sapiens

125 &lt;400&gt; SEQUENCE: 2

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127 1 5 10 15
128 Thr Glu Arg Ser Phe Pro Arg Ala Ser Glu Asp Glu Val Thr Cys Pro
129 20 25 30
130 Thr Ser Ala Pro Pro Ser Pro Thr Arg Thr Arg Gly Asn Cys Ala Glu
131 35 40 45
132 Ala Glu Glu Gly Gly Cys Arg Gly Ala Pro Arg Lys Leu Arg Ala Arg
133 50 55 60
134 Arg Gly Gly Arg Ser Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
135 65 70 75 80
136 Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
137 85 90 95
138 His Asn Leu Asn Ser Ala Leu Asp Ala Leu Arg Gly Val Leu Pro Thr
139 100 105 110
140 Phe Pro Asp Asp Ala Lys Leu Thr Lys Ile Glu Thr Leu Arg Phe Ala
141 115 120 125
142 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
143 130 135 140
144 Ser Leu Tyr Ala Leu Glu Pro Pro Ala Pro His Cys Gly Glu Leu Gly
145 145 150 155 160
146 Ser Pro Gly Gly Ser Pro Gly Asp Trp Gly Ser Leu Tyr Ser Pro Val
147 165 170 175
148 Ser Gln Ala Gly Ser Leu Ser Pro Ala Ala Ser Leu Glu Glu Arg Pro
149 180 185 190
150 Gly Leu Leu Gly Ala Thr Ser Ser Ala Cys Leu Ser Pro Gly Ser Leu
151 195 200 205
152 Ala Phe Ser Asp Phe Leu
153 210
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156 &lt;211&gt; LENGTH: 1861

157 &lt;212&gt; TYPE: DNA

158 &lt;213&gt; ORGANISM: Mus musculus

160 &lt;400&gt; SEQUENCE: 3

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163 ggttgacacac acatttcctg acccaaccca tagtggcgga gagctggata gcactttgaa 180
164 ctaatgggcyg ctcctcccag ctgccagcca agaagacact tgactccttg atcgctggtt 240
165 catttagaca agccgtttcc ctctctgagc caaaagaccc catgtgtaat actcaaagaa 300
166 gaggccttcc ttatatatat ataggcacc ccaaacctcc ttcattgctac caagaaaggg 360
167 tctggacaca tgccaaaaag aaagaggaaa aggcacagct ctcccagcg gccggacggg 420
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169 cgggggaggg gtggagtggg ggaacaaaca gactgctgct cccctcccc gaccctgcc 540
170 ctttgtccgg aactccagct tgctctgcgg gtgggggttg tggggggagg agcgggctcg 600
171 cgtggcgcaag cccctgggcc cccctccgctg attggcccg gtgcaggca gcagcccgcc 660
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179 cccaaccgca ggatggcgcc tcctcccttg gatgcgtca ccatccaagt gtccccagag 1140
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190 gacctgtctg gctctgggtg gtgggtgcta gtggaaagg aggggaccag agccgtctgg 1800
191 agtgggaggt agtggaggct ctcaagcacc tcgcctcttc tggctttcac tacttggtac 1860
192 c 1861
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194 &lt;210&gt; SEQ ID NO: 4

195 &lt;211&gt; LENGTH: 214

196 &lt;212&gt; TYPE: PRT

197 &lt;213&gt; ORGANISM: Mus musculus

199 &lt;400&gt; SEQUENCE: 4

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201 1 5 10 15
202 Thr Gln Gln Pro Phe Pro Gly Ala Ser Asp His Glu Val Leu Ser Ser
203 20 25 30
204 Asn Ser Thr Pro Pro Ser Pro Thr Leu Ile Pro Arg Asp Cys Ser Glu
205 35 40 45
206 Ala Glu Val Gly Asp Cys Arg Gly Thr Ser Arg Lys Leu Arg Ala Arg
207 50 55 60
208 Arg Gly Gly Arg Asn Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
209 65 70 75 80
210 Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
211 85 90 95
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216 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
217           130           135           140
218 Ser Phe Tyr Gly Pro Glu Pro Pro Val Pro Cys Gly Glu Leu Gly Ser
219 145           150           155           160
220 Pro Gly Gly Gly Ser Asn Gly Asp Trp Gly Ser Ile Tyr Ser Pro Val
221           165           170           175
222 Ser Gln Ala Gly Asn Leu Ser Pro Thr Ala Ser Leu Glu Glu Phe Pro
223           180           185           190
224 Gly Leu Gln Val Pro Ser Ser Pro Ser Tyr Leu Leu Pro Gly Ala Leu
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232 <213> ORGANISM: Artificial Sequence
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235 <223> OTHER INFORMATION: oligonucleotide primer
237 <400> SEQUENCE: 5
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240 <210> SEQ ID NO: 6
241 <211> LENGTH: 23
242 <212> TYPE: DNA
243 <213> ORGANISM: Artificial Sequence
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246 <223> OTHER INFORMATION: primer
248 <400> SEQUENCE: 6
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252 <211> LENGTH: 16
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial Sequence
256 <220> FEATURE:
257 <223> OTHER INFORMATION: oligonucleotide primer
259 <400> SEQUENCE: 7
260 atcctgcggt tgggga
262 <210> SEQ ID NO: 8
263 <211> LENGTH: 21
264 <212> TYPE: DNA
265 <213> ORGANISM: Artificial Sequence
267 <220> FEATURE:
268 <223> OTHER INFORMATION: oligonucleotide primer
270 <400> SEQUENCE: 8
271 tggaagggtgt gtgtgtgccca g
273 <210> SEQ ID NO: 9

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VERIFICATION SUMMARY

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Input Set : A:\SeqList for UCSF-129CIP.txt

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L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date